AMENDMENT

Please amend the claims without prejudice, without admission, without surrender of the subject matter, and without any intention of creating any estoppel as to equivalents as follows.

IN THE CLAIMS

Please amend the claims as follows:

23. (Thrice Amended) A method for identifying elite event MS-B2 in a transgenic *Brassica* plant, or cell or tissue thereof, or transgenic *Brassica* plant material, which method comprises detecting in the DNA of said plant, cell, tissue or material a MS-B2 specific region with a specific primer or probe which hybridizes to bases 1-234 of SEQ ID NO:8 or to bases 194-416 of SEQ ID NO:10 of MS-B2.

E1

24. (Thrice Amended) The method of claim 23, said method comprising amplifying a DNA fragment of between 160 and 200 by from a nucleic acid present in said transgenic *Brassica* plant, or cell or tissue thereof, or transgenic *Brassica* plant material, using a polymerase chain reaction with at least two primers, one of which hybridizes to bases 1-234 of SEQ ID NO:8 or to bases 194-416 of SEQ ID NO:10 of MS-B2, the other of which hybridizes to a sequence within SEQ ID NO:1, and detecting said amplified DNA fragment on an agarose gel.

25. (Amended) The method of claim 24, wherein one of said primers hybridizes to a sequence within SEQ ID NO:1 and comprises the sequence of SEQ ID NO:12.

E2

26. (Amended) The method of claim 24, wherein one of said primers hybridizes to bases 194-416 of SEQ ID NO:10 and comprises the sequence of SEQ ID NO:11.

29. (Twice Amended) A kit for identifying elite event MS-B2 in a transgenic *Brassica* plant, or cell or tissue thereof, or transgenic *Brassica* plant material, said kit comprising at least one PCR primer, which hybridizes to bases 1-234 of SEQ ID NO:8 or to bases 194-416 of SEQ ID NO:10 of MS-B2.

E3

30. (Amended) The kit of Claim 29, which further comprises at least a second PCR primer which hybridizes to a sequence within SEQ ID NO:1 of MS-B2.

E4

31. (Amended) The kit of claim 30, wherein said PCR primer comprises the sequence of SEQ ID NO:12.

32. (Amended) The kit of claim 29, wherein said at least one PCR primer comprises the sequence of SEQ ID NO:11.

34. (Twice Amended) A method for screening the genomic DNA of seeds for the presence of MS-B2, which method comprises detecting an MS-B2 specific region comprising the insertion site of MS-B2 with a specific primer or probe which hybridizes to bases 1-234 of SEQ ID NO:8 or to bases 194-416 of SEQ ID NO:10 of MS-B2, and thus confirming the presence of MS-B2 if the MS-B2 specific DNA sequence is so detected in samples of seed lots.

35. (Amended) A method for screening the genomic DNA of seeds for the absence of MS-B2, which method comprises carrying out a Polymerase Chain Reaction or Southern Blot using a specific primer or probe which hybridizes to bases 1-234 of SEQ ID NO:8 or to bases 194-416 of SEQ ID NO:10 of MS-B2, and not detecting the presence of MS-B2 specific DNA on an agarose gel or Southern Blot membrane, thus confirming the absence of MS-B2 in said seeds.

36. (Amended) A method for identifying a *Brassica* plant, or cell or tissue thereof, or *Brassica* plant material not comprising elite event MS-B2, which method comprises performing a polymerase chain reaction with at least two primers, one of which recognizes bases 1-234 of SEQ ID NO:8 or bases 194-416 of SEQ ID NO:10 of MS-B2, another of which recognizes a sequence within SEQ ID NO:1, and detecting the absence of a DNA fragment of between 160 and 200 base pairs on an agarose gel.

Please cancel claims 27, 28 and 33 without prejudice.